## **CLAIMS**

Therefore, having thus described the disclosure, at least the following is claimed:

- 1 An isolated polynucleotide comprising a polynucleotide selected from: a polynucleotide
- 2 sequence set forth in SEQ ID NO: 1 (C307YhGALE) or a degenerate variant of the SEQ ID
- NO: 1; a polynucleotide sequence at least 90% identical to the polynucleotide sequence set forth
- 4 in SEQ ID NO: 1; a polynucleotide sequence at least 75% identical to the polynucleotide
- 5 sequence set forth in SEQ ID NO: 1; and a polynucleotide sequence at least 50% identical to the
- 6 polynucleotide sequence set forth in SEQ ID NO: 1.
- 1 2. A polypeptide selected from: an amino acid sequence set forth in SEQ ID NO: 2
- 2 (C307YhGALE), or conservatively modified variants thereof; an amino acid sequence that is at
- least 90% identical to SEQ ID NO: 2; an amino acid sequence that is at least 75% identical to
- 4 SEQ ID NO: 2; and an amino acid sequence that is at least 50% identical to SEQ ID NO: 2.
- 1 3. A vector comprising the isolated polynucleotide of claim 1.
- 1 . 4. The vector of claim 3 wherein the vector is pPIC3.5K.
- 1 5. An isolated host cell comprising the vector of claim 3.
- 1 6. The isolated host cell of claim 5 wherein the host cell is selected from: *Pichia pastoris*,
- 2 Saccharomyces cerevisiae, Schizosaccharomyces pombe, and Escherichia coli.
- The isolated host cell of claim 6 wherein the host cell is *Pichia pastoris*.

- 8. A process for producing a polypeptide comprising culturing the host cell of claim 7 under
- 2 conditions sufficient for the production of the polypeptide where the polypeptide has the
- 3 characteristics that the polypeptide is capable of UDP-gal/UDP-glc interconversion and
- 4 substantially incapable of UDP-galNAc/UDP-glcNAc interconversion.
- 1 9. The process of claim 8 wherein the polypeptide is the polypeptide of claim 2.
- 1 10. A cell line transfected with an expression vector comprising a polynucleotide selected
- 2 from: a polynucleotide sequence set forth in SEQ ID NO: 1(C307YhGALE) or a degenerate
- 3 variant of the SEQ ID NO: 1; a polynucleotide sequence at least 90% identical to the
- 4 polynucleotide sequence set forth in SEQ ID NO: 1; a polynucleotide sequence at least 75%
- 5 identical to the polynucleotide sequence set forth in SEQ ID NO: 1; and a polynucleotide
- sequence at least 50% identical to the polynucleotide sequence set forth in SEQ ID NO: 1,
- 7 encoding a polypeptide having the characteristics that the polypeptide is capable of UDP-
- 8 gal/UDP-glc interconversion and substantially incapable of UDP-galNAc/UDP-glcNAc
- 9 interconversion.
- 1 11. The cell line of claim 10 wherein the polypeptide is selected from: an amino acid
- 2 sequence set forth in SEQ ID NO: 2 (C307YhGALE), or conservatively modified variants
- 3 thereof; an amino acid sequence that is at least 90% identical to SEQ ID NO: 2; an amino acid
- 4 sequence that is at least 75% identical to SEQ ID NO: 2; and an amino acid sequence that is at
- 5 least 50% identical to SEQ ID NO: 2.
- 1 12. The cell line of claim 10 wherein the expression vector is pCDNA3.
- 1 13. The cell line of claim 10 wherein the cell line is GALE deficient.

- 1 14. The cell line of claim 13 wherein the cell line is *ldlD*.
- 1 15. A vector comprising an isolated polynucleotide selected from: a polynucleotide sequence
- 2 set forth in SEQ ID NO: 3 (WTeGALE), or a degenerate variant of the SEQ ID NO: 3; a
- polynucleotide sequence at least 90% identical to the polynucleotide sequence set forth in SEQ
- 4 ID NO: 3; a polynucleotide sequence at least 75% identical to the polynucleotide sequence set
- forth in SEQ ID NO: 3; and a polynucleotide sequence at least 50% identical to the
- 6 polynucleotide sequence set forth in SEQ ID NO: 3.
- 1 16. The vector of claim 15 wherein the vector is pPIC3.5K.
- 1 17. An isolated host cell comprising the vector of claim 15.
- 1 18. The isolated host cell of claim 17 wherein the host cell is selected from: *Pichia pastoris*,
- 2 Saccharomyces cerevisiae, Schizosaccharomyces pombe, and Escherichia coli.
- 1 The isolated host cell of claim 18 wherein the host cell is *Pichia pastoris*.
- 1 20. A process for producing a polypeptide comprising culturing the host cell of claim 19
- 2 under conditions sufficient for the production of the polypeptide where the polypeptide has the
- 3 characteristics that the polypeptide is capable of UDP-gal/UDP-glc interconversion and
- 4 substantially incapable of UDP-galNAc/UDP-glcNAc interconversion.
- 1 21. The process of claim 20 wherein the polypeptide is selected from: an amino acid
- 2 sequence set forth in SEQ ID NO: 4, or conservatively modified variants thereof; an amino acid
- 3 sequence that is at least 90% identical to SEQ ID NO: 4; an amino acid sequence that is at least

- 4 75% identical to SEQ ID NO: 4; and an amino acid sequence that is at least 50% identical to
- 5 SEQ ID NO: 4
- 1 22. A cell line transfected with an expression vector comprising a polynucleotide selected
- from: a polynucleotide SEQ ID NO: 3 (WTeGALE) or a degenerate variant of the SEQ ID NO:
- 3; a polynucleotide sequence at least 90% identical to the polynucleotide sequence set forth in
- 4 SEQ ID NO: 3; a polynucleotide sequence at least 75% identical to the polynucleotide sequence
- set forth in SEQ ID NO: 3; and a polynucleotide sequence at least 50% identical to the
- 6 polynucleotide sequence set forth in SEQ ID NO: 3 encoding a polypeptide having the
- 7 characteristics that the polypeptide is capable of UDP-gal/UDP-glc interconversion and
- 8 substantially incapable of UDP-galNAc/UDP-glcNAc interconversion.
- 1 23. The cell line of claim 22 wherein the polypeptide is selected from: an amino acid
- 2 sequence set forth in SEQ ID NO: 4 (WTeGALE), or conservatively modified variants thereof;
- an amino acid sequence that is at least 90% identical to SEQ ID NO: 4; an amino acid sequence
- 4 that is at least 75% identical to SEQ ID NO: 4; and an amino acid sequence that is at least 50%
- 5 identical to SEQ ID NO: 4
- 1 24. The cell line of claim 22 wherein the expression vector is pCDNA3.
- 1 25. The cell line of claim 22 wherein the cell line is GALE deficient.
- 1 26. The cell line of claim 25 wherein the cell line is *ldlD*.
- 1 27. A method of culturing the cell line of claim 10 in the absence of galactose to produce
- 2 glycoproteins having N-linked modifications with substantially no O-linked modifications.

- 1 28. A method of culturing the cell line of claim 22 in the absence of galactose to produce
- 2 glycoproteins having N-linked modifications with substantially no O-linked modifications.